SEQUENCE LISTING

<110> Giesing, Michael
Suchy, Bernhard

<120> METHOD FOR ANALYZING BODY FLUIDS FOR THE PRESENCE OF CANCER CELLS, USE THEREOF, CORRESPONDING ANALYSIS KITS, AND USE OF SPECIFIC ACTIVE SUBSTANCES FOR TREATING CANCER

<130> GIES3002

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<151> 2003-08-20

<150> DE 102 38 046.5
<151> 2002-08-20

<160> 22

<170> PatentIn version 3.3

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<220>
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<210> 13
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20 25 30

Tyr Asp Tyr Gly Ala Leu Glu Pro His Ile Asn Ala Gln Ile Met Gln
35 40 45

Leu His His Ser Lys His His Ala Ala Tyr Val Asn Asn Leu Asn Val
50 55 60

Thr Glu Glu Lys Tyr Gln Glu Ala Leu Ala Lys Gly Asp Val Thr Ala
65 70 75 80

Gln Thr Ala Leu Gln Pro Ala Leu Lys Phe Asn Gly Gly Gly His Ile
85 90 95

Asn His Ser Ile Phe Trp Thr Asn Leu Ser Pro Asn Gly Gly Gly Glu
100 105 110

Pro Lys Gly Glu Leu Leu Glu Ala Ile Lys Arg Asp Phe Gly Ser Phe

115 120 125

Asp Lys Phe Lys Glu Lys Leu Thr Ala Ala Ser Val Gly Val Gln Gly
130 135 140

Ser Gly Trp Gly Trp Leu Gly Phe Asn Lys Glu Arg Gly His Leu Gln
145 150 155 160

Ile Ala Ala Cys Pro Asn Gln Asp Pro Leu Gln Gly Thr Thr Gly Leu
165 170 175

Ile Pro Leu Leu Gly Ile Asp Val Trp Glu His Ala Tyr Tyr Leu Gln
180 185 190

Tyr Lys Asn Val Arg Pro Asp Tyr Leu Lys Ala Ile Trp Asn Val Ile
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Asn Trp Glu Asn Val Thr Glu Arg Tyr Met Ala Cys Lys Lys
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<213> Homo sapiens

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Gln Tyr Gly Lys Lys Val Met Val Leu Asp Phe Val Thr Pro Thr Pro
        35                  40                  45

Leu Gly Thr Arg Trp Gly Leu Gly Gly Thr Cys Val Asn Val Gly Cys
    50                  55                  60

Ile Pro Lys Lys Leu Met His Gln Ala Ala Leu Leu Gly Gln Ala Leu
65                  70                  75                  80

Gln Asp Ser Arg Asn Tyr Gly Trp Lys Val Glu Glu Thr Val Lys His
                85                  90                  95

Asp Trp Asp Arg Met Ile Glu Ala Val Gln Asn His Ile Gly Ser Leu
            100                 105                 110

Asn Trp Gly Tyr Arg Val Ala Leu Arg Glu Lys Lys Val Val Tyr Glu
        115                 120                 125

Asn Ala Tyr Gly Gln Phe Ile Gly Pro His Arg Ile Lys Ala Thr Asn
    130                 135                 140

Asn Lys Gly Lys Glu Lys Ile Tyr Ser Ala Glu Ser Phe Leu Ile Ala
145                 150                 155                 160

Thr Gly Glu Arg Pro Arg Tyr Leu Gly Ile Pro Gly Asp Lys Glu Tyr
                165                 170                 175

Cys Ile Ser Ser Asp Asp Leu Phe Ser Leu Pro Tyr Cys Pro Gly Lys
            180                 185                 190

Thr Leu Val Val Gly Ala Ser Tyr Val Ala Leu Glu Cys Ala Gly Phe
        195                 200                 205

Leu Ala Gly Ile Gly Leu Gly Val Thr Val Met Val Arg Ser Ile Leu
    210                 215                 220

Leu Arg Gly Phe Asp Gln Asp Met Ala Asn Lys Ile Gly Glu His Met
225                 230                 235                 240

Glu Glu His Gly Ile Lys Phe Ile Arg Gln Phe Val Pro Ile Lys Val
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245 250 255

Glu Gln Ile Glu Ala Gly Thr Pro Gly Arg Leu Arg Val Val Ala Gln
260 265 270

Ser Thr Asn Ser Glu Glu Ile Ile Glu Gly Glu Tyr Asn Thr Val Met
275 280 285

Leu Ala Ile Gly Arg Asp Ala Cys Thr Arg Lys Ile Gly Leu Glu Thr
290 295 300

Val Gly Val Lys Ile Asn Glu Lys Thr Gly Lys Ile Pro Val Thr Asp
305 310 315 320

Glu Glu Gln Thr Asn Val Pro Tyr Ile Tyr Ala Ile Gly Asp Ile Leu
325 330 335

Glu Asp Lys Val Glu Leu Thr Pro Val Ala Ile Gln Ala Gly Arg Leu
340 345 350

Leu Ala Gln Arg Leu Tyr Ala Gly Ser Thr Val Lys Cys Asp Tyr Glu
355 360 365

Asn Val Pro Thr Thr Val Phe Thr Pro Leu Glu Tyr Gly Ala Cys Gly
370 375 380

Leu Ser Glu Glu Lys Ala Val Glu Lys Phe Gly Glu Glu Asn Ile Glu
385 390 395 400

Val Tyr His Ser Tyr Phe Trp Pro Leu Glu Trp Thr Ile Pro Ser Arg
405 410 415

Asp Asn Asn Lys Cys Tyr Ala Lys Ile Ile Cys Asn Thr Lys Asp Asn
420 425 430

Glu Arg Val Val Gly Phe His Val Leu Gly Pro Asn Ala Gly Glu Val
435 440 445

Thr Gln Gly Phe Ala Ala Ala Leu Lys Cys Gly Leu Thr Lys Lys Gln
450 455 460

Leu Asp Ser Thr Ile Gly Ile His Pro Val Cys Ala Glu Val Phe Thr
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Thr Leu Ser Val Thr Lys Arg Ser Gly Ala Ser Ile Leu Gln Ala Gly
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Cys

<210> 16

<211> 1314
<212> DNA
<213> Homo sapiens

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cggcctgccg gcggggacga cagcattgcg cctgggtgca gcagtgtgcg tctcggggaa 180

gggaagatat tttaaggcgt gtctgagcag acggggaggc ttttccaaac ccaggcagct 240

tcgtggcgtg tgcggtttcg acccggtcac acaaagcttc agcatgtcat gtgaggacgg 300

tcgggccctg aaaggaacgc tctcggaatt ggccgcggaa accgatctgc ccgttgtgtt 360

tgtgaaacag agaaagatag gcggccatgg tccaaccttg aaggcttatc aggagggcag 420

acttcaaaag ctactaaaaa tgaacggccc tgaagatctt cccaagtcct atgactatga 480

ccttatcatc attggaggtg gctcaggagg tctggcagct gctaaggagg cagcccaata 540

tggcaagaag gtgatggtcc tggactttgt cactcccacc cctcttggaa ctagatgggg 600

tcttggagga acatgtgtga atgtgggttg catacctaaa aaactgatgc atcaagcagc 660

tttgttagga caagccctgc aagactctcg aaattatgga tggaaagtcg aggagacagt 720

taagcatgat tgggacagaa tgatagaagc tgtacagaat cacattggct ctttgaattg 780

gggctaccga gtagctctgc gggagaaaaa agtcgtctat gagaatgctt atgggcaatt 840

tattggtcct cacaggatta aggcaacaaa taataaaggc aaagaaaaaa tttattcagc 900

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agaatactgc atcagcagtg atgatctttt ctccttgcct tactgcccgg gtaagaccct 1020

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aggcgtcact gttatggtta ggtccattct tcttagagga tttgaccagg acatggccaa 1140

caaaattggt gaacacatgg aagaacatgg catcaagttt ataagacagt tcgtaccaat 1200

taaagttgaa caaattgaag cagggacacc aggccgactc agagtagtag ctcagtccac 1260

caatagtgag gaaatcattg aaggagaata taatacggtg atgctggcaa tagg 1314

<210> 17
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<213> Homo sapiens

<400> 17

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Phe Ser Ala Arg Pro Leu Ala Gly Gly Glu Pro Val Ser Leu Gly Ser
20 25 30

Leu Arg Gly Lys Val Leu Leu Ile Glu Asn Val Ala Ser Leu Cys Gly
35 40 45

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Thr Thr Val Arg Asp Tyr Thr Gln Met Asn Glu Leu Gln Arg Arg Leu
    50                  55                  60

Gly Pro Arg Gly Leu Val Val Leu Gly Phe Pro Cys Asn Gln Phe Gly
65                  70                  75                  80

His Gln Glu Asn Ala Lys Asn Glu Glu Ile Leu Asn Ser Leu Lys Tyr
                85                  90                  95

Val Arg Pro Gly Gly Gly Phe Glu Pro Asn Phe Met Leu Phe Glu Lys
            100                 105                 110

Cys Glu Val Asn Gly Ala Gly Ala His Pro Leu Phe Ala Phe Leu Arg
        115                 120                 125

Glu Ala Leu Pro Ala Pro Ser Asp Asp Ala Thr Ala Leu Met Thr Asp
    130                 135                 140

Pro Lys Leu Ile Thr Trp Ser Pro Val Cys Arg Asn Asp Val Ala Trp
145                 150                 155                 160

Asn Phe Glu Lys Phe Leu Val Gly Pro Asp Gly Val Pro Leu Arg Arg
                165                 170                 175

Tyr Ser Arg Arg Phe Gln Thr Ile Asp Ile Glu Pro Asp Ile Glu Ala
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Leu Leu Ser Gln Gly Pro Ser Cys Ala
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<213> Homo sapiens

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gaggcaccac ggtccgggac tacacccaga tgaacgagct gcagcggcgc ctcggacccc    240

ggggcctggt ggtgctcggc ttcccgtgca accagtttgg gcatcaggag aacgccaaga    300

acgaagagat tctgaattcc ctcaagtacg tccggcctgg tggtgggttc gagcccaact    360

tcatgctctt cgagaagtgc gaggtgaacg gtgcgggggc gcaccctctc ttcgccttcc    420

tgcgggaggc cctgccagct cccagcgacg acgccaccgc gcttatgacc gaccccaagc    480

tcatcacctg gtctccggtg tgtcgcaacg atgttgcctg gaactttgag aagttcctgg    540

tgggccctga cggtgtgccc ctacgcaggt acagccgccg cttccagacc attgacatcg    600

agcctgacat cgaagccctg ctgtctcaag ggcccagctg tgcctagggc gcccctccta    660
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agtgccgggt gtcagc 856

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<212> DNA
<213> Artificial

<220>
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<210> 20
<211> 60
<212> DNA
<213> Artificial

<220>
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<210> 21
<211> 60
<212> DNA
<213> Artificial

<220>
<223> probe (GPX2)

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<210> 22
<211> 60
<212> DNA
<213> Artificial

<220>
<223> probe (GPX3)

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